

GenCore version 5.1.4-p5-4578
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OM protein - protein search, using sw model

Run on: March 20, 2003, 03:52:32 ; Search time 63 Seconds

(without alignments)
329.954 Million cell updates/sec

Title: US-09-867-958-1

Perfect score: 849
Sequence: 1 MARQHAKRLMTDRPMYVME.....PPAMDLDSDSDADATSN 156

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database :

A.Geneseq.101002:*

- 1: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*
- 2: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
- 3: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1982.DAT:*
- 4: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1983.DAT:*
- 5: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1984.DAT:*
- 6: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1985.DAT:*
- 7: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1986.DAT:*
- 8: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:*
- 9: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1988.DAT:*
- 10: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1989.DAT:*
- 11: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1990.DAT:*
- 12: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1991.DAT:*
- 13: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1992.DAT:*
- 14: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
- 15: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1994.DAT:*
- 16: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1995.DAT:*
- 17: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:*
- 18: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1997.DAT:*
- 19: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1998.DAT:*
- 20: /SID52/gcgdata/geneseq/geneseqp-emb1/AA1999.DAT:*
- 21: /SID52/gcgdata/geneseq/geneseqp-emb1/AA2000.DAT:*
- 22: /SID52/gcgdata/geneseq/geneseqp-emb1/AA2001.DAT:*
- 23: /SID52/gcgdata/geneseq/geneseqp-emb1/AA2002.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	ID	Description
1	849	100.0	156 20 AAY02591	A human progesterone
2	843	99.3	156 22 AAM39556	Human polypeptide
3	772	90.9	543 22 AAM39557	Human polypeptide
4	771	90.8	619 22 AAM82707	Human immune/haema
5	690	81.3	196 22 AAM41342	Human polypeptide
6	347.5	40.9	160 22 AAG63379	Amino acid sequenc
7	347.5	40.9	180 23 ABR41768	Human ovarian anti
8	238	28.0	239 22 AAU30460	Novel human secret
9	126	14.8	69 21 AAB59081	Breast and ovarian
10	126	14.8	134 22 AAO02613	Human polypeptide

11	125	14.7	371	22 ABB64334	Drosophila melanog
12	121	14.3	244	21 AAG09812	Arabidopsis thalia
13	121	14.3	258	21 AAG47700	Arabidopsis thalia
14	121	14.3	262	21 AAG32123	Arabidopsis thalia
15	115	13.5	362	22 AAM93819	Human polypeptide
16	115	13.5	362	22 AAM38809	Human polypeptide
17	115	13.5	362	22 AAB94033	Human protein sequ
18	115	13.5	371	21 AAB56812	Human prostate can
19	115	13.5	371	22 AAM40695	Human polypeptide
20	115	13.5	371	22 AAM40696	Human polypeptide
21	91.5	10.8	704	9 AAP80087	Sequence of 85 kd
22	91.5	10.8	704	23 AAM71813	Trypanosoma anti
23	89.5	10.5	327	12 AAM14026	Human brain tissu
24	89.5	10.5	487	18 AAM26510	Amyloid precursor
25	89.5	10.5	487	18 AAM26394	Amyloid precursor
26	89.5	10.5	487	19 AAM44745	APP-REP 751 protel
27	89.5	10.5	487	19 AAM42978	Amyloid precursor
28	89.5	10.5	492	14 AAM45229	APP-REP 751 amylo
29	89.5	10.5	492	18 AAM26509	Amyloid precursor
30	89.5	10.5	492	18 AAM26393	APP-REP 751 prc
31	89.5	10.5	492	19 AAM44744	Amyloid precursor
32	89.5	10.5	492	19 AAM42978	Beta-amyloid precu
33	89.5	10.5	596	15 AAM65797	Sequence of amylo
34	89.5	10.5	651	15 AAM65796	APP695. Homo sapl
35	89.5	10.5	670	15 AAM65795	APP695 mutant A-de
36	89.5	10.5	695	9 AAP81692	APP695 mutant A-de
37	89.5	10.5	695	12 AAR05166	APP695 mutant A-de
38	89.5	10.5	695	11 AAR14046	APP695 mutant A-de
39	89.5	10.5	695	13 AAR26338	APP695 mutant A-de
40	89.5	10.5	695	18 AAM19487	APP695 mutant A-de
41	89.5	10.5	695	18 AAM19501	APP695 mutant A-de
42	89.5	10.5	695	18 AAM19504	APP695 mutant A-de
43	89.5	10.5	695	18 AAM19498	APP695 mutant A-de
44	89.5	10.5	695	18 AAM19484	APP695 mutant A-de
45	89.5	10.5	695	18 AAM19495	APP695 mutant A-de

ALIGNMENTS

RESULT 1	AA02591	standard; Protein; 156 AA.
ID	AA02591	
XX	AA02591;	
AC	19-JUL-1999	(first entry)
XX		
XX		A human progesterone receptor complex p23-like protein.
DE		
XX		Human progesterone receptor complex p23-like protein; PR23P;
KW		neurological disorder; antagonist; reproductive disorder;
KW		immunological disorder; neoplastic disorder.
XX		
OS	Homo sapiens	
XX		
PN	MO9919483-AL	
XX	22-APR-1999	
PD	09-OCT-1998;	98WO-US21402.
XX	09-OCT-1997;	97US-0948197.
PR		
XX		
PA	(INCY-) INCYTE PHARM INC.	
XX		
PI	Corley NC, Shah P, Yue H;	
XX		
DR	WPI: 1999-302530/25.	
XX	N-PSDB; AAX36136.	
PT		
XX		Human progesterone receptor complex p23-like protein

PS Claim 1; Fig 1A-B; 67pp; English.

XX

CC The present sequence represents a human progesterone receptor complex

CC p23-like protein (PR23p). PR23p is used to treat neurological

CC disorders. Antagonists of PR23p are useful for treating reproductive,

CC immunological or neoplastic disorders. Probes and primers based on the

CC PR23p polynucleotides can be used for diagnosis, detection and screening

CC of homologues, and amplification of PR23p genes. Antisense PR23p

CC polynucleotides can be used to decrease or inhibit expression of PR23p.

XX

SO Sequence 156 AA;

Query Match 100.0%; Score 849; DB 20; Length 156;

Best Local Similarity 100.0%; Pred. No. 1e-82;

Matches 156; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAROHARTLWTDPRMYVMEFCVEDSTVHVLIEDHRIYFSCKNADGYELNIEIFYAKV 60

DB 1 MAROHARTLWTDPRMYVMEFCVEDSTVHVLIEDHRIYFSCKNADGYELNIEIFYAKV 60

QY 61 NSKSDKRRSRSTICFVRKKKEKVAWPLRTKEDIKPYWLSYDFDNNWRDWEDEEMELAH 120

DB 61 NSKSDKRRSRSTICFVRKKKEKVAWPLRTKEDIKPYWLSYDFDNNWRDWEDEEMELAH 120

QY 121 VEHYAELLKKVSTKRPPAMDLDSDSADDAATSN 156

DB 121 VEHYAELLKKVSTKRPPAMDLDSDSADDAATSN 156

RESULT 2

AAAM39556

XX AAAM39556; Protein: 156 AA.

XX

AC AAAM39556;

XX

DT 22-OCT-2001 (first entry)

XX

DE Human polypeptide SEQ ID NO 2701.

XX

XX Human; noctropic; immunosuppressant; cytostatic; gene therapy; cancer;

XX Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic;

XX amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;

XX chemokinetic; thrombolytic; drug screening; arthritis; inflammation;

XX leukaemia.

XX

OS Homo sapiens.

XX

PN WO20015312-A1.

XX

PD 26-JUL-2001.

XX

PF 26-DEC-2000; 2000WO-US34263.

XX

PR 21-JAN-2000; 2000US-0488725.

PR 25-APR-2000; 2000US-0552317.

PR 09-JUL-2000; 2000US-0598042.

PR 19-JUL-2000; 2000US-0620312.

PR 03-AUG-2000; 2000US-0653450.

PR 14-SEP-2000; 2000US-0662191.

PR 19-OCT-2000; 2000US-0693036.

PR 29-NOV-2000; 2000US-0727344.

XX

PA (HYSE-) HYSEQ INC.

XX

PI Tang YT, Liu C, Asundi V, Chen R, Ma Y, Qian XB, Ren F, Wang D;

PI Wang J, Wang Z, Wehrman T, Xu C, Xue AJ, Yang Y, Zhang J;

PI Zhao QA, Zhou P, Goodrich R, Drmanac RT;

XX

DR WPI; 2001-442253/47.

DR N-PSDB; AAI58712.

XX

PT Novel nucleic acids and polypeptides, useful for treating disorders

PT such as central nervous system injuries -

XX

PS Example 4; SEQ ID NO 2701; 10078pp; English.

XX

CC The invention relates to human nucleic acids (AAI57798-AAI61369) and

CC the encoded polypeptides (AAAM38642-AAAM42213) with noctropic,

CC immunosuppressant and cytostatic activity. The polynucleotides are useful

CC in gene therapy. A composition containing a polypeptide or polynucleotide

CC of the invention may be used to treat diseases of the peripheral nervous

CC system, such as peripheral nervous injuries, peripheral neuropathy and

CC localized neuropathies and central nervous system diseases, such as

CC Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic

CC lateral sclerosis, and Shy-Drager Syndrome. Other uses include the

CC utilization of the activities such as: Immune system suppression,

CC Activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic

CC and thrombolytic activity, cancer diagnosis and therapy, drug screening,

CC assays for receptor activity, arthritis and inflammation, leukaemias and

CC C.N.S disorders.

CC Note: The sequence data for this patent did not form part of the prin

CC specification.

XX

SO Sequence 156 AA;

Query Match 99.3%; Score 843; DB 22; Length 156;

Best Local Similarity 99.4%; Pred. No. 4.5e-82;

Matches 155; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAROHARTLWTDPRMYVMEFCVEDSTVHVLIEDHRIYFSCKNADGYELNIEIFYAKV 60

DB 1 MAROHARTLWTDPRMYVMEFCVEDSTVHVLIEDHRIYFSCKNADGYELNIEIFYAKV 60

QY 61 NSKSDKRRSRSTICFVRKKKEKVAWPLRTKEDIKPYWLSYDFDNNWRDWEDEEMELAH 120

DB 61 NSKSDKRRSRSTICFVRKKKEKVAWPLRTKEDIKPYWLSYDFDNNWRDWEDEEMELAH 120

QY 121 VEHYAELLKKVSTKRPPAMDLDSDSADDAATSN 156

DB 121 VEHYAELLKKVSTKRPPAMDLDSDSADDAATSN 156

RESULT 3

AAAM39657

XX AAAM39657; Protein: 543 AA.

XX

AC AAAM39657;

XX

DT 22-OCT-2001 (first entry)

XX

DE Human polypeptide SEQ ID NO 2802.

XX

XX Human; noctropic; immunosuppressant; cytostatic; gene therapy; cancer;

XX Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic;

XX amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;

XX chemokinetic; thrombolytic; drug screening; arthritis; inflammation;

XX leukaemia.

XX

OS Homo sapiens.

XX

PN WO20015312-A1.

XX

PD 26-JUL-2001.

XX

PF 26-DEC-2000; 2000WO-US34263.

XX

PR 21-JAN-2000; 2000US-0488725.

PR 25-APR-2000; 2000US-0552317.

PR 09-JUL-2000; 2000US-0598042.

PR 19-JUL-2000; 2000US-0620312.

PR 03-AUG-2000; 2000US-0653450.

PR 14-SEP-2000; 2000US-0662191.

PR 19-OCT-2000; 2000US-0693036.

PR 29-NOV-2000; 2000US-0727344.

PS	Claim 11; SEQ ID NO 10300; 3071pp + Sequence Listing; English.
XX	
CC	AAK54951 to AAK64702 encode the human immune/hematopoietic antigen (I)
CC	amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic
CC	activity, and can be used in gene therapy and vaccine production. (I)
CC	proteins and polynucleotides may be used in the prevention, diagnosis and
CC	treatment of diseases associated with inappropriate (I) expression. For
CC	example, they may be used to treat disorders associated with decreased
CC	expression by rectifying mutations or deletions in a patient's genome
CC	that affect the activity of (I) by expressing inactive proteins or to
CC	supplement the patients own production of (I). Additionally, (I)
CC	polynucleotides may be used to produce the secreted (I), by inserting
CC	the nucleic acids into a host cell and culturing the cell to express the
CC	protein. (II) proteins and polynucleotides may be used to prevent,
CC	diagnose and treat immune/haematopoietic-related diseases, especially
CC	cancers and cancer metastases of haematopoietic-derived cells. AAK64703
CC	to AAK87654 represent human immune/haematopoietic antigen genomic
CC	sequences from the present invention. AAK54942 to AAK54950 and AAM821f
CC	represent sequences used in the exemplification of the present inventi
XX	
SO	Sequence 619 AA;
	Query Match 90.8%; Score 771; DB 22; Length 619;
	Best Local Similarity 97.9%; Pred. No. 1.3e-73;
	Matches 141; Conservative 0; Mismatches 3; Indels 0; Gaps 0
OY	1 MAROHARLWMDRPKYVMEFCVEEDSTVHVLIEDHRIYFSCKNADGVELNYIEFYAKV 60
Db	61 MARKHARLWMDRPRYVMEFCVEDSTVHVLIEDHRIYFSCKNADGVELNYIEFYAKX 120
OY	61 NSKSQDSRRSRJITCFVRKKMKEXAMPRLTRKEDIKPWLSDPDMNRMDEGEDELNAH 120
Db	121 NSKSQDSRRSRJITCFVRKKMKEXAMPRLTREDIKPWLSDPDMNRMDEGEDELNAH 180
OY	121 VEHYAELLKKVSTRKPPAMDLD 144
Db	181 VEHYAELLKKVSTRKPPAMDLD 204
RESULT 5	
AAAM1342	
ID	AAM41342 standard; Protein; 196 AA.
AC	AAM41342;
XX	
DT	22-OCT-2001 (first entry)
XX	
DE	Human polypeptide SEQ ID NO 6273.
KW	Human; noctropic; immunosuppressant; cytosstatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Draeger Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening; arthritis; inflammation; leukemia.
OS	Homo sapiens.
XX	
PB	MO20015312-A1.
PD	26-JUL-2001.
PF	26-DEC-2000; 200OWO-US34263.
XX	
PR	21-JAN-2000; 200OUS-0488725.
PR	25-APR-2000; 200OUS-0552317.
PR	09-JUL-2000; 200OUS-0598042.
PR	19-JUL-2000; 200OUS-0620312.
PR	03-AUG-2000; 200OUS-0653450.
PR	14-SEP-2000; 200OUS-0662191.
PR	19-OCT-2000; 200OUS-0693036.
PR	29-NOV-2000; 200OMS-0727344.

XX (HYSE-) HYSEQ INC.
 PA
 XX Tang YF, Liu C, Asundi V, Chen R, Ma Y, Qian XB, Ren F, Wang D;
 PI Wang J, Wang Z, Weinman T, Xu C, Xue AJ, Yang Y, Zhang J;
 PI Zhao QA, Zhou P, Goodrich R, Drmanac RT;
 XX
 DR WPI: 2001-442253/47.
 DR N-PsDB: AA160498.
 XX
 PT Novel nucleic acids and polypeptides, useful for treating disorders
 PT such as central nervous system injuries -
 XX
 PS Example 2; SEQ ID NO 6273; 10078pp; English.
 XX
 CC The invention relates to human nucleic acids (AA157798-AA161369) and
 CC the encoded polypeptides (AA158642-AA162213) with nootropic,
 CC immunosuppressant and cytostatic activity. The polynucleotides are useful
 CC in gene therapy. A composition containing a polypeptide or polynucleotide
 CC of the invention may be used to treat diseases of the peripheral nervous
 CC system, such as peripheral nervous injuries, peripheral neuropathy and
 CC localised neuropathies and central nervous system diseases, such as
 CC Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic
 CC lateral sclerosis, and Shy-Drager Syndrome. Other uses include the
 CC utilisation of the activities such as: Immune system suppression,
 CC Activin/Inhibin activity, chemotactic/chemokinetic activity, haemostatic
 CC and thrombolytic activity, cancer diagnosis and therapy, drug screening,
 CC assays for receptor activity, arthritis and inflammation, leukaemias and
 CC C.N.S. disorders.
 CC Note: The sequence data for this patent did not form part of the printed
 CC specification.
 CC
 XX
 SQ Sequence 196 AA;
 Query Match 81.3%; Score 690; DB 22; Length 196;
 Best Local Similarity 98.4%; Pred. No. 1.3e-65;
 Matches 125; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
 QY 1 MARQHARTLWYDRPMYVEMFECVEDSTDVHVLIEDHRIYFSCKNADGVELYNEIEFYAKV 60
 DB 60 MARQHARTLWYDRPMYVEMFECVEDSTDVHVLIEDHRIYFSCKNADGVELYNEIEFYAKV 119
 QY 61 NSKDSQDKRSSSTICFVAKKMEKYAMPRLTKEDIKPVMLSYDFDWMRDWEGDEEMELAH 120
 DB 120 NSKDSQDKRSSSTICFVAKKMEKYAMPRLTKEDIKPVMLSYDFDWMRDWEGDEEMELAH 179
 QY 121 VEHYAEI. 127
 DB 180 VEHYAEV 186

RESULT 6
 AAG63379
 ID AAG63379 standard; Protein, 160 AA.
 XX
 AC AAG63379;
 XX
 DT 15-OCT-2001 (first entry)
 DE Amino acid sequence of a human prostaglandin E1 (PGE1) synthase.
 XX
 KW Human; prostaglandin E1 synthase; PGE1 synthase; arachidonic acid;
 KW inflammation.
 XX
 OS Homo sapiens.
 OS
 PN WO200157225-A1.
 XX
 PD 09-AUG-2001.
 XX
 PF 25-AUG-2000; 2000WO-JP05758.
 XX
 PR 03-FEB-2000; 2000JP-0032704.

XX (CHUS) CHUGAI SEIYAKU KK.
 PA (KUDO/) KUDO I.
 XX
 PI Kudo I, Murakami M, Ohishi S;
 XX
 DR WPI: 2001-483439/52.
 DR N-PsDB: AA43100.
 XX
 PT PGE2-1 protein and encoded gene with PGE2 synthase activity, useful in
 PT screening efficient PGE2 synthase inhibitors as antiinflammatory agents
 PT
 XX
 PS Claim 1; Fig 5; 54pp; Japanese.
 XX
 CC The present sequence represents a human prostaglandin E1 (PGE1) synthase.
 CC The protein synthesizes PGE2 from arachidonic acid in consort with COX.
 CC The PGE2 synthase protein and gene are useful in screening for efficient
 CC PGE2 synthase inhibitors. These inhibitors are useful as
 CC anti-inflammatory agents.
 CC
 XX
 SQ Sequence 160 AA;
 Query Match 40.9%; Score 347.5; DB 22; Length 160;
 Best Local Similarity 43.4%; Pred. No. 3.9e-29;
 Matches 66; Conservative 30; Mismatches 53; Indels 3; Gaps 2;
 QY 4 QAHARTLWYDRPMYVEMFECVEDSTDVHVLIEDHRIYFSC-KNADGVELYNEIEFYAKVNS 62
 DB 2 QPASAKMYDRDQVYFTEFCVEDSKDVANFEKSKTFSCIGSDNFKHLNEIDLFHCIDP 61
 QY 63 KDSQDKRSSSTICFVAKKMEKYAMPRLTKEDIKPVMLSYDFDWMRDWEGDEEMELAHVE 122
 DB 62 NDSKKHRTDRSILCCLRKSGSQWPRLTKEKAKLNLVDFENKWKWEDSDSDMSNFD 121
 QY 123 HYAEILKVVSTKRPP--PAMDDIDDDSDSADD 152
 DB 122 RFSEMMNNMGDEDVDLPFYDAGADDDSDSDSD 153

RESULT 7
 ABP41768
 ID ABP41768 standard; Protein, 180 AA.
 XX
 AC ABP41768;
 XX
 DT 22-AUG-2002 (first entry)
 DE Human ovarian antigen HOPK759, SEQ ID NO:2900.
 XX
 KW Human; ovarian antigen; ovary; ovarian; breast; cancer; tumour;
 KW ovarian cancer; breast cancer; tumour; reproductive system disorder;
 KW infertility; pregnancy disorder; anovulation; polycystic ovary syndrome;
 KW PCOS; ovarian cyst; dysmenorrhoea; endocrine disorder; infection;
 KW inflammatory condition; immune disorder; blood disorder;
 KW cardiovascular disorder; respiratory disorder; neurological disorder;
 KW gastrointestinal disorder; urinary system disorder; drug screening;
 KW gene therapy; chromosome mapping; forensic analysis;
 KW antibody preparation; cytostatic; immunomodulatory; neuroprotective;
 KW antiinflammatory; gynaecological; reproductive.
 XX
 OS Homo sapiens.
 OS
 PN WO200200677-A1.
 XX
 PD 03-JAN-2002.
 XX
 PF 07-JUN-2001; 2001WO-US18569.
 XX
 PR 07-JUN-2000; 2000US-209467P.
 XX
 PA (HUMA-) HUMAN GENOME SCI INC.

PI Birse CE, Rosen CA;
XX
DR WPI: 2002-147878/19.
DR N-PSDB: ABQ54845.
XX
PT Isolated nucleic acid molecules encoding novel ovarian polypeptides,
PT useful in the prevention, treatment and diagnosis of cancer (e.g.
PT ovarian cancer), immune disorders, cardiovascular disorders and
PT neurological diseases -
XX
PS Claim 11; SEQ ID No 2900; 2922pp; English.
XX
CC The invention relates to 2175 novel human ovarian antigens (ABP41054-
CC ABP43228) and to cDNAs encoding them (ABQ54131-ABQ56305), and also
CC encompasses polypeptides 90% identical and polynucleotides 95% identical
CC to the sequences of the invention. The invention additionally relates to
CC recombinant vectors and host cells comprising human ovarian antigen
CC polynucleotides, antibodies against human ovarian antigens, and the use
CC of ovarian antigen polynucleotides and polypeptides in diagnosing,
CC treating, prognosing or preventing various ovary and/or breast-related
CC disorders. Such conditions include ovarian cancer and breast cancer, and
CC metastatic tumours of ovarian or breast origin, reproductive system
CC disorders (e.g., infertility, disorders of pregnancy, anovulation,
CC polycystic ovary syndrome, ovarian cysts, and dysmenorrhoea), endocrine
CC disorders, infections (e.g., chlamydia, HIV, toxoplasmosis, and toxic
CC shock syndrome), inflammatory conditions (e.g., mastitis, oophoritis and
CC vaginitis), immune disorders (e.g., congenital and acquired
CC immunodeficiencies, autoimmune oophoritis, systemic lupus erythematosus),
CC blood-related disorders (e.g., anaemia), cardiovascular disorders,
CC respiratory disorders, neurological disorders, gastrointestinal disorders
CC and urinary system disorders. Ovarian antigen polypeptides and
CC polynucleotides may also be used in screening for compounds which
CC modulate ovarian antigen expression or activity. The polynucleotides may
CC further be used for gene therapy, chromosome mapping, in the
CC identification of individuals and in forensic analysis, and the
CC polypeptides may be used as food additives or to prepare antibodies
CC useful in disease diagnosis, drug targeting and phenotyping. The present
CC sequence represents a human ovarian antigen of the invention.
CC Note: The sequence data for this patent did not form part of the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 180 AA;
Query Match 40.9%; Score 347.5; DB 23; Length 180;
Best Local Similarity 43.4%; Pred. No. 4.5e-29;
Matches 66; Conservative 30; Mismatches 53; Indels 3; Gaps 2;
OY 4 QHARTLWDRPMYVMECEVEDSTDVHLLIEDHRIYFSC-KNADGVELYNEIEFPAKVS 62
DB 22 QPASKAKWYDRDYPVIEFCVEDSKDVNVNFEKSKLTFSCLGSDNFKHNEIDLFHCIDP 81
OY 63 KDSODKRSSRSITCFVRKMKKEKVPAMRLTKEDIKFPWLSVDGNRDMRGDEMEELAHVE 122
DB 82 NDSKHKRRDRSILCLCKRGESGSPRLTKERAKLWLSVDENNMKDWEDSDDEDSNFD 141
OY 123 HYAELLKRVSTKRPP--PAMDLDLDDSDSADD 152
DB 142 RFSEMMNNMGDEVDLPEYDGDADDSDSD 173

RESULT 8
AAU30460
ID AAU30460 standard; Protein: 239 AA.
XX
AC AAU30460;
XX
DT 18-DEC-2001 (first entry)
XX
DE Novel human secreted protein #951.
XX
DE Human; vaccination; gene therapy; nutritional supplement;
KM stem cell proliferation; haematopoiesis; nerve tissue regeneration;

KW Immune suppression; immune stimulation; anti-inflammatory; leukaemia.
XX
OS Homo sapiens.
XX
PN WO200179449-A2.
XX
PN 25-OCT-2001.
PD
XX
PF 16-APR-2001; 2001WO-US08656.
XX
PR 18-APR-2000; 2000US-0552929.
PR 26-JAN-2001; 2001US-0770160.
XX
PA (HYSE-) HYSEQ INC.
XX
PI Tang YF, Liu C, Drmanac RT;
XX
DR WPI: 2001-611725/70.
XX
PT Nucleic acids encoding a range of human polypeptides, useful in genet.
PT vaccination, testing and therapy -
XX
PS Claim 20; Page 296; 765pp; English.
XX
CC The invention relates to novel human secreted polypeptides. The
CC polypeptides and antibodies to the polypeptides are useful for
CC determining the presence of or predisposition to a disease associated
CC with altered levels of polypeptide. The polypeptides are also useful for
CC identifying agents (agonists and antagonists) that bind to them. Cells
CC expressing the proteins are useful for identifying a therapeutic agent
CC for use in treatment of a pathology related to aberrant expression or
CC physiological interactions of the polypeptide. Vectors comprising
CC the nucleic acids encoding the polypeptides and cells genetically
CC engineered to express them are also useful for producing the proteins.
CC The proteins are useful in genetic vaccination, testing and
CC therapy, and can be used as nutritional supplements. They may be used to
CC increase stem cell proliferation; to regulate haematopoiesis; and in
CC bone, cartilage, tendon and/or nerve tissue growth or regeneration;
CC immune suppression and/or stimulation; as anti-inflammatory agents; and
CC in treatment of leukemias. AAU29510-AAU3304 represent the amino acid
CC sequences of novel human secreted proteins of the invention.
XX
SQ Sequence 239 AA;
Query Match 28.0%; Score 238; DB 22; Length 239;
Best Local Similarity 36.8%; Pred. No. 3.3e-17;
Matches 60; Conservative 30; Mismatches 55; Indels 10; Gaps 1;
OY 4 QHARTLWDRPMYV-MEFC-VEDSTDVHLLIEDHRIY-FSC---KNADGVELYNEIEF 56
DB 74 QPASKAKWYDRDYPVIEFCVEDSKDVNVNFEKSKLTFSCLGSDNFKHNEIDLFHCIDP 131
OY 57 YAKVNSKDSODKRSSRSITCFVRK--WKKEKAMPRLTKEDIKFPWLSVDGNRDMRGDE 111
DB 132 FHCIDPNDSKHKRRDTSILCLCKRGESGSPRLTKERAKLWLSVDENNMKDWEDSDDE 189
OY 112 GDEMEELAHVEHYAELLKRVSTKR--PPAMDLDLDDSDSADD 152
DB 190 DSDDEMSNFDKRFSEMMNNMGDEVDYDPEYDGDADDSDSD 232

RESULT 9
AAB59081
ID AAB59081 standard; Protein: 69 AA.
XX
AC AAB59081;
XX
DT 27-MAR-2001 (first entry)
XX
DE Breast and ovarian cancer associated antigen protein sequence SPQ ID 789.
XX
DE Human; breast cancer; ovarian cancer; cytostatic; immunosuppressive;
KM neoplastic; neuroprotective; antiviral; antiallergic; hepatotropic;

XX	Human polypeptide SEQ ID NO 16505.
DE	
XX	Human; cytokine; cell proliferation; cell differentiation; gene therapy;
KW	vaccine; peptide therapy; stem cell growth factor; haematopoiesis;
KW	tissue growth factor; immunomodulatory; cancer; leukaemia;
KW	nervous system disorders; arthritis; inflammation.
XX	
OS	Homo sapiens.
XX	
PN	WO200164835-A2.
XX	
PD	07-SEP-2001.
XX	
PF	26-FEB-2001; 2001WO-US04927.
XX	
PR	28-FEB-2000; 2000US-0515126.
PR	18-MAY-2000; 2000US-0577409.
XX	
PA	(HYSE-) HYSEQ INC.
XX	
PI	Tang YT, Liu C, Dymanac RT;
XX	
XX	WPI: 2001-514838/56.
DR	N-PSDB; AA182544.
DR	
XX	
PT	Isolated nucleic acids and polypeptides, useful for preventing
PT	diagnosing and treating e.g. leukaemia, inflammation and immune
PT	disorders -
XX	
ES	Claim 20; SEQ ID NO 16505; 1399bp + Sequence Listing; English.
XX	
XX	The invention relates to human polynucleotides (AA179941-AA193841) and
CC	the encoded proteins (AA000010-AA013910) that exhibit activity elating to
CC	cytokine, cell proliferation or cell differentiation or which may induce
CC	production of other cytokines in other cell populations. The
CC	polynucleotides and polypeptides are useful in gene therapy, vaccines or
CC	peptide therapy. The polypeptides have various cytokine-like activities,
CC	e.g. stem cell growth factor activity, haematopoiesis regulating
CC	activity, tissue growth factor activity, immunomodulatory activity and
CC	activin/inhibin activity and may be useful in the diagnosis and/or
CC	treatment of cancer, leukaemia, nervous system disorders, arthritis and
CC	inflammation.
CC	Note: The sequence data for this patent did not form part of the printed
CC	specification, but was obtained in electronic format directly from WIPO
CC	at ftp.wipo.int/pub/published_pct_sequences.
XX	
XX	
SO	Sequence 134 AA;
	Query Match 14.8%; Score 126; DB 22; Length 134;
	Best Local Similarity 24.8%; Pred. No. 1.4e-05;
	Matches 31; Conservative 31; Mismatches 49; Indels 14; Gaps 3
OY	19 MEFCEVDSTDVHVLIEDHRIYVSCKNADGVELYNLEFPAKYNKSQDKRRSRSTICFV 78
DB	8 LNLCTTASLKHFFSIARKF-----HLNGSHLQSO-HSCEAQLRRTARPIICL 56
OY	79 RKMKEKVAWPRLETKDIPKWLSDVDFDMNRDWDGDEMELAHNEHVAELIKKYSTKRP 138
DB	57 PYGEGGQSMPIITLERANINLWLTVFNNMKDWEYDSQYMSNDRSEEMNIIIVMKR--- 113
OY	139 AMDL 143
DB	114 SIDNL 118
	RESULT 11
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ID	ABB64334 standard; Protein; 371 AA.
XX	
XX	ABB64334;
XX	
CT	26-MAR-2002 (first entry)

XX DE Drosophila melanogaster polypeptide SEQ ID NO 19794.
XX KW Drosophila; developmental biology; cell signalling; insecticide;
XX KW pharmaceutical.
OS Drosophila melanogaster.
XX PN WO200171042-A2.
XX PD 27-SEP-2001.
XX PF 23-MAR-2001; 2001WO-US09231.
XX PR 23-MAR-2000; 2000US-191637P.
XX PR 11-JUL-2000; 2000US-0614150.
XX PA (PEKE) PE CORP NY.
XX PI Venter JC, Adams M, Li PWD, Myers EW;
XX DR WPI: 2001-656860/75.
XX DR N-PSDB: ABL08437.
XX PT New isolated nucleic acid detection reagent for detecting 1000 or more
XX PT genes from Drosophila and for elucidating cell signalling and cell-cell
XX PT interactions -
XX PS Disclosure; SEQ ID NO 19794; 21pp + Sequence Listing; English.
XX XX
XX CC The invention relates to an isolated nucleic acid detection reagent
XX CC capable of detecting 1000 or more genes from Drosophila. The invention is
XX CC useful in developmental biology and in elucidating cell signalling and
XX CC cell-cell interactions in higher eukaryotes for the development of
XX CC insecticides, therapeutics and pharmaceutical drugs. The invention
XX CC discloses genomic DNA sequences (AB16176-ABL30511), expressed DNA
XX CC sequences (AB161840-ABL16175) and the encoded proteins
XX CC (ABB57737-ABB72072).
XX CC The sequence data for this patent did not form part of the printed
XX CC specification, but was obtained in electronic format directly from WIPO
XX CC at ftp.wipo.int/pub/published_pct_sequences.
XX CC
SQ Sequence 371 AA;
Query Match 14.7%; Score 125; DB 22; Length 371;
Best Local Similarity 36.2%; Pred. NO. 6.7e-05;
Matches 34; Conservative 13; Mismatches 37; Indels 10; Gaps 4;
OY 45 ADGVELYN-EIEFYAKVNSKDSODKRSRSTCEFRKKKEKVAAPRLTKEDIKPYWLSVD 103
DB 47 ARGVNAVKFELHFXALIDENATFVSDNKTLEQIRK-LPEPMWRPLVATPOKPHMLKID 105
OY 104 FDNWRDMEGDEME-----LAHYEYAEILKK 130
DB 106 FDRWRT-EDDVEVEKEPRDVQDEKEXADLOKR 138
RESULT 12
AAG09812
ID AAG09812 standard; Protein; 244 AA.
XX
XX AAG09812;
XX
XX 17-OCT-2000 (first entry)
XX
XX Arabidopsis thaliana protein fragment SEQ ID NO: 7889.
XX
XX Protein identification; signal transduction pathway; metabolic pathway;
XX KW hybridisation assay; genetic mapping; gene expression control; promoter;
XX KW termination sequence.
XX Arabidopsis thaliana.
XX

PN EP1033405-A2.
XX
XX PD 06-SEP-2000.
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XX PF 25-FEB-2000; 2000EP-0301439.
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XX PF 25-FEB-1999; 99US-0121825.
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XX EPI033405-A2.
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Query Match 14.3%; Score 121; DB 21; Length 258;
Best Local Similarity 27.6%; Pred. No. 0.00011;

Matches 35; Conservative 25; Mismatches 57; Indels 10; Gaps 5;

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OY 62 SKDSQDKRSSRITCFVRKKKEKVPRLTKEDIKPYWLSVDFDNMDWEGDE-----E 115
DB 63 VESKINIGERSIFCJIEK-AEPEERNKKLIRVKKPPHYAVVDMKWVD-EDDEGSAGAD 120
OY 116 MELAHVE 122
DB 121 MDWAGME 127

RESULT 14
AAG32123
ID AAG32123 standard; Proteln; 262 AA.

XX AAG32123;

DT 17-OCT-2000 (first entry)

DE Arabidopsis thaliana protein fragment SEQ ID NO: 38695.

XX Protein identification; signal transduction pathway; metabolic pathway;
KW hybridisation assay; genetic mapping; gene expression control; promoter;
XX termination sequence.

XX

OS Arabidopsis thaliana.
XX
XX EPI033405-A2.
XX
XX 06-SEP-2000.
XX
XX 25-FEB-2000; 2000EP-0301439.
XX
PR 25-FEB-1999; 990S-0121825.
PR 05-MAR-1999; 990S-0123180.
PR 09-MAR-1999; 990S-0123548.
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DB 125 MDMAQWE 131  
  
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XX Human; full length cDNA; cDNA synthesis; oligo-capping.  
OS Homo sapiens.
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PR 02-MAY-2000; 2000JP-0183765.
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PA (HELI-) HELIX RES INST.
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PI Ota T, Nishikawa T, Isogai T, Hayashi K, Ishii S, Kawai Y;
PI Wakamatsu A, Sugiyama T, Nagai K, Kojima S, Otsuki T, Koga H;
XX
DR MPI; 2001-524255/58.
DR N-PSDB; AAK94775.
XX
PT 830 Primers useful for synthesizing full length cDNA clones and their
PT use in genetic manipulation.
XX
PS Claim 8; SEQ ID NO 3873; 1380bp + sequence listing; English.
XX
CC The invention relates to primers for synthesizing full length cDNA
CC clones. 830 cDNA molecules encoding a human protein have been
CC isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA
CC molecules have been determined. Primers for synthesizing the full length
CC cDNA are useful for clarifying the function of the protein encoded by
CC the cDNA. The full length clones were obtained by construction of full
CC length enriched cDNA libraries that were synthesised by the oligo-capping
CC method. The primers enable the production of the full length cDNA easily
CC without any special methods. The present sequence is a polypeptide
CC Note: The sequence data for this patent did not form part of the printed
CC specification, but was obtained in CD-ROM format directly from EPO.
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SQ Sequence 362 AA;

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Query Match 13.5%; Score 115; DB 22; Length 362;
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